

each position along the sequences divided by the total length of the aligned sequences.

Gaps are considered non-identical residues for the purpose of the calculations.

Applicants respectfully request that the executed Declaration and remarks be entered and made of record in the instant application.

Date: May 25, 2001

Respectfully submitted,

*By* *Adriane M. Antler* *Reg. No. 31,232*  
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